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IN THE UNITED STATES PATENT AND TRADEMARK OFFICE

APPLICANTS: Willson et al.
SERIAL NO.: 09/994,701
FILED: 01/17/01

§ ART UNIT NO.: 1651
§ EXAMINER: UNKNOWN
§ DOCKET NO.: 96605/13UTL
§

TITLE: Nucleic Acid Separation Using
Immobilized Metal Affinity Chromatography

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Robert W. Strozier	Washington, D.C. 20231	June 24, 2002
		Date of Signature

STATEMENT REGARDING SUBSTITUTE PAPER COPY OF SEQUENCE LISTING AND CRF COPY OF SEQUENCE LISTING

Dear Sir/Madam:

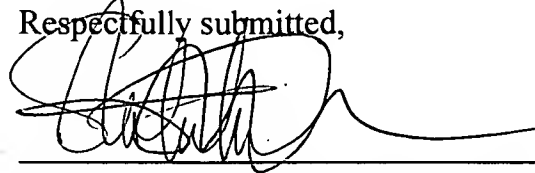
In response to a Notice to Comply with Sequence Rules, Applicant used the PatentIn 3.1 software for the United States Patent and Trademark Office to generate a hard copy and electronic copy of the sequence listing as required by the Notice.

Applicants' Attorney verifies that the paper and electronic listing are identical, both generated by PatentIn 3.1.

Applicants' Attorney has attempted in good faith to respond to this notice; however, Applicants' Attorney must point out that at numerous part of the specification, reference is made to specific DNA or RNA samples. These reference are to particular plasmids or other nucleic acid sequence, without the actual sequence. Applicants did not include the actual sequences of these nucleic acids because the actual sequences are no germane to this application. The application separations nucleic acid sequences based on base richness.

If this submission is considered deficient, please have someone from the sequence compliance group call me. I have attempted several times to get an answer on the issue of whether a nucleic acid sequence listing is required for a mere reference to a specific nucleic acid sequence, without sequence listing, in an application.

Respectfully submitted,



Date: June 24, 2002

Robert W. Strozier, Reg. No. 34,024
Attorney for Applicants